

ma
tran

1642

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/462,625

ENTERED

CRF Processing Date: 5/25/2001

Edited by: [Signature]
Verified by: [Signature] (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☒ Other: Seq. 2 - added 42207

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/462,625

DATE: 05/25/2001

TIME: 14:38:22

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05252001\I462625.raw

4 <110> APPLICANT: Georgiev, Georgii
5 Kiselev, Sergei
6 Prokhorchouk, Egor
7 Ostermann, Elinborg
9 <120> TITLE OF INVENTION: Tumor Growth Inhibition- and Apoptosis-Associated Genes
10 and Polypeptides and Methods of Use Thereof
12 <130> FILE REFERENCE: 0652.1630001
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/462,625
C--> 15 <141> CURRENT FILING DATE: 2000-07-28
17 <150> PRIOR APPLICATION NUMBER: US 08/893,764
18 <151> PRIOR FILING DATE: 1997-07-11
20 <150> PRIOR APPLICATION NUMBER: PCT/EP98/04287
21 <151> PRIOR FILING DATE: 1998-07-10
23 <160> NUMBER OF SEQ ID NOS: 24
25 <170> SOFTWARE: PatentIn Ver. 2.1
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28 <211> LENGTH: 549
29 <212> TYPE: DNA
30 <213> ORGANISM: Artificial Sequence
32 <220> FEATURE:
33 <221> NAME/KEY: CDS
34 <222> LOCATION: (1)..(549)
36 <220> FEATURE:
37 <223> OTHER INFORMATION: Description of Artificial Sequence: tag7 cDNA
39 <400> SEQUENCE: 1
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42 1 5 10 15
44 tgc agt ttc atc gtg ccc cgc agt gag tgg agg gcc ctg cca tcc gag 96
45 Cys Ser Phe Ile Val Pro Arg Ser Glu Trp Arg Ala Leu Pro Ser Glu
46 20 25 30
48 tgc tct agc cgc ctg ggg cac cca gtt cgc tac gtg gtg atc tca cac 144
49 Cys Ser Ser Arg Leu Gly His Pro Val Arg Tyr Val Val Ile Ser His
50 35 40 45
52 aca gcc ggc agc ttc tgc aac agc ccg gac tcc tgt gaa cag cag gcc 192
53 Thr Ala Gly Ser Phe Cys Asn Ser Pro Asp Ser Cys Glu Gln Gln Ala
54 50 55 60
56 cgc aat gtg cag cat tac cac aag aat gag ctg ggc tgg tgc gat gta 240
57 Arg Asn Val Gln His Tyr His Lys Asn Glu Leu Gly Trp Cys Asp Val
58 65 70 75 80
60 gcc tac aac ttc ctt att gga gag gac ggt cat gtc tat gaa ggc cga 288
61 Ala Tyr Asn Phe Leu Ile Gly Glu Asp Gly His Val Tyr Glu Gly Arg
62 85 90 95
64 ggc tgg aac atc aag ggt gac cac aca ggg ccc atc tgg aat ccc atg 336
65 Gly Trp Asn Ile Lys Gly Asp His Thr Gly Pro Ile Trp Asn Pro Met
66 100 105 110
68 tct att ggc atc acc ttc atg ggg aac ttc atg gac cgg gta ccc gca 384

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Input Set : A:\Pto.amc

Output Set: C:\CRF3\05252001\I462625.raw

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70      115      120      125
72 aag cgg gcc ctc cgt gct gcc cta aat ctt ctg gaa tgt ggg gtg tct 432
73 Lys Arg Ala Leu Arg Ala Ala Leu Asn Leu Leu Glu Cys Gly Val Ser
74      130      135      140
76 cgg ggc ttc ctg aga tcc aac tat gaa gtc aaa gga cac cgg gat gtg 480
77 Arg Gly Phe Leu Arg Ser Asn Tyr Glu Val Lys Gly His Arg Asp Val
78 145      150      155      160
80 caa agc act ctc tct cca ggt gac caa ctc tat cag gtc atc caa agc 528
81 Gln Ser Thr Leu Ser Pro Gly Asp Gln Leu Tyr Gln Val Ile Gln Ser
82      165      170      175
84 tgg gaa cac tac cga gag tga 549
85 Trp Glu His Tyr Arg Glu
86      180
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90 <211> LENGTH: 182
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94 <220> FEATURE:
95 <223> OTHER INFORMATION: Description of Artificial Sequence: tag7 cDNA
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100 Cys Ser Phe Ile Val Pro Arg Ser Glu Trp Arg Ala Leu Pro Ser Glu
101 20 25 30
102 Cys Ser Ser Arg Leu Gly His Pro Val Arg Tyr Val Val Ile Ser His
103 35 40 45
104 Thr Ala Gly Ser Phe Cys Asn Ser Pro Asp Ser Cys Glu Gln Gln Ala
105 50 55 60
106 Arg Asn Val Gln His Tyr His Lys Asn Glu Leu Gly Trp Cys Asp Val
107 65 70 75 80
108 Ala Tyr Asn Phe Leu Ile Gly Glu Asp Gly His Val Tyr Glu Gly Arg
109 85 90 95
110 Gly Trp Asn Ile Lys Gly Asp His Thr Gly Pro Ile Trp Asn Pro Met
111 100 105 110
112 Ser Ile Gly Ile Thr Phe Met Gly Asn Phe Met Asp Arg Val Pro Ala
113 115 120 125
114 Lys Arg Ala Leu Arg Ala Ala Leu Asn Leu Leu Glu Cys Gly Val Ser
115 130 135 140
116 Arg Gly Phe Leu Arg Ser Asn Tyr Glu Val Lys Gly His Arg Asp Val
117 145 150 155 160
118 Gln Ser Thr Leu Ser Pro Gly Asp Gln Leu Tyr Gln Val Ile Gln Ser
119 165 170 175
120 Trp Glu His Tyr Arg Glu
121 180
125 <210> SEQ ID NO: 3
126 <211> LENGTH: 718
127 <212> TYPE: DNA
128 <213> ORGANISM: Homo sapiens

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/462,625

DATE: 05/25/2001

TIME: 14:38:22

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05252001\I462625.raw

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131 <221> NAME/KEY: 5'UTR
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134 <220> FEATURE:
135 <221> NAME/KEY: CDS
136 <222> LOCATION: (68)..(643)
138 <220> FEATURE:
139 <221> NAME/KEY: 3'UTR
140 <222> LOCATION: (644)..(718)
142 <220> FEATURE:
143 <221> NAME/KEY: polyA_site
144 <222> LOCATION: (712)..(714)
146 <400> SEQUENCE: 3
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149 ccgctct atg ctg ctt gcc tgg gct ctc ccc agc ctc ctt cga ctc gga 109
150 Met Leu Leu Ala Trp Ala Leu Pro Ser Leu Leu Arg Leu Gly
151 1 5 10
153 gcg gct cag gag aca gaa gac ccg gcc tgc tgc agc ccc ata gtg ccc 157
154 Ala Ala Gln Glu Thr Glu Asp Pro Ala Cys Cys Ser Pro Ile Val Pro
155 15 20 25 30
157 cgg aac gag tgg aag gcc ctg gca tca gag tgc gcc cag cac ctg agc 205
158 Arg Asn Glu Trp Lys Ala Leu Ala Ser Glu Cys Ala Gln His Leu Ser
159 35 40 45
161 ctg ccc tta cgc tat gtg gtg gta tcg cac acg gcg ggc agc agc tgc 253
162 Leu Pro Leu Arg Tyr Val Val Val Ser His Thr Ala Gly Ser Ser Cys
163 50 55 60
165 aac acc ccc gcc tcg tgc cag cag cag gcc cgg aat gtg cag cac tac 301
166 Asn Thr Pro Ala Ser Cys Gln Gln Gln Ala Arg Asn Val Gln His Tyr
167 65 70 75
169 cac atg aag aca ctg ggc tgg tgc gac gtg ggc tac aac ttc ctg att 349
170 His Met Lys Thr Leu Gly Trp Cys Asp Val Gly Tyr Asn Phe Leu Ile
171 80 85 90
173 gga gaa gac ggg ctc gta tac gag ggc cgt ggc tgg aac ttc acg ggt 397
174 Gly Glu Asp Gly Leu Val Tyr Glu Gly Arg Gly Trp Asn Phe Thr Gly
175 95 100 105 110
177 gcc cac tca ggt cac tta tgg aac ccc atg tcc att ggc atc agc ttc 445
178 Ala His Ser Gly His Leu Trp Asn Pro Met Ser Ile Gly Ile Ser Phe
179 115 120 125
181 atg ggc aac tac atg gat cgg gtg ccc aca ccc cag gcc atc cgg gca 493
182 Met Gly Asn Tyr Met Asp Arg Val Pro Thr Pro Gln Ala Ile Arg Ala
183 130 135 140
185 gcc cag ggt cta ctg gcc tgc ggt gtg gct cag gga gcc ctg agg tcc 541
186 Ala Gln Gly Leu Leu Ala Cys Gly Val Ala Gln Gly Ala Leu Arg Ser
187 145 150 155
189 aac tat gtg ctc aaa gga cac cgg gat gtg cag cgt aca ctc tct cca 589
190 Asn Tyr Val Leu Lys Gly His Arg Asp Val Gln Arg Thr Leu Ser Pro
191 160 165 170
193 ggc aac cag ctc tac cac ctc atc cag aat tgg cca cac tac cgc tcc 637
194 Gly Asn Gln Leu Tyr His Leu Ile Gln Asn Trp Pro His Tyr Arg Ser

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/462,625

DATE: 05/25/2001

TIME: 14:38:22

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05252001\I462625.raw

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195 175                      180                      185                      190
197 ccc tga ggccctgctg atccgcaccc cattcctccc ctcccatggc caaaaacccc      693
198 Pro
200 actgtctcct tctccaataa agatg      718
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204 <211> LENGTH: 191
205 <212> TYPE: PRT
206 <213> ORGANISM: Homo sapiens
208 <400> SEQUENCE: 4
209 Met Leu Leu Ala Trp Ala Leu Pro Ser Leu Leu Arg Leu Gly Ala Ala
210 1 5 10 15
211 Gln Glu Thr Glu Asp Pro Ala Cys Cys Ser Pro Ile Val Pro Arg Asn
212 20 25 30
213 Glu Trp Lys Ala Leu Ala Ser Glu Cys Ala Gln His Leu Ser Leu Pro
214 35 40 45
215 Leu Arg Tyr Val Val Val Ser His Thr Ala Gly Ser Ser Cys Asn Thr
216 50 55 60
217 Pro Ala Ser Cys Gln Gln Gln Ala Arg Asn Val Gln His Tyr His Met
218 65 70 75 80
219 Lys Thr Leu Gly Trp Cys Asp Val Gly Tyr Asn Phe Leu Ile Gly Glu
220 85 90 95
221 Asp Gly Leu Val Tyr Glu Gly Arg Gly Trp Asn Phe Thr Gly Ala His
222 100 105 110
223 Ser Gly His Leu Trp Asn Pro Met Ser Ile Gly Ile Ser Phe Met Gly
224 115 120 125
225 Asn Tyr Met Asp Arg Val Pro Thr Pro Gln Ala Ile Arg Ala Ala Gln
226 130 135 140
227 Gly Leu Leu Ala Cys Gly Val Ala Gln Gly Ala Leu Arg Ser Asn Tyr
228 145 150 155 160
229 Val Leu Lys Gly His Arg Asp Val Gln Arg Thr Leu Ser Pro Gly Asn
230 165 170 175
231 Gln Leu Tyr His Leu Ile Gln Asn Trp Pro His Tyr Arg Ser Pro
232 180 185 190
236 <210> SEQ ID NO: 5
237 <211> LENGTH: 14
238 <212> TYPE: DNA
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
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245 tttttttttt ttac      14
248 <210> SEQ ID NO: 6
249 <211> LENGTH: 10
250 <212> TYPE: DNA
251 <213> ORGANISM: Artificial Sequence
253 <220> FEATURE:
254 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
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RAW SEQUENCE LISTING

DATE: 05/25/2001

PATENT APPLICATION: US/09/462,625

TIME: 14:38:22

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05252001\I462625.raw

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262 <212> TYPE: DNA
263 <213> ORGANISM: Artificial Sequence
265 <220> FEATURE:
266 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
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273 <211> LENGTH: 18
274 <212> TYPE: DNA
275 <213> ORGANISM: Artificial Sequence
277 <220> FEATURE:
278 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
280 <400> SEQUENCE: 8
281 gaaagggcct cgtgatac                             18
284 <210> SEQ ID NO: 9
285 <211> LENGTH: 36
286 <212> TYPE: DNA
287 <213> ORGANISM: Artificial Sequence
289 <220> FEATURE:
290 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
292 <400> SEQUENCE: 9
293 cagggccctc cactcacttg cataggcatt tgtagc         36
296 <210> SEQ ID NO: 10
297 <211> LENGTH: 36
298 <212> TYPE: DNA
299 <213> ORGANISM: Artificial Sequence
301 <220> FEATURE:
302 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
304 <400> SEQUENCE: 10
305 gctacaaatg cctatgcaag tgagtggagg gccctg         36
308 <210> SEQ ID NO: 11
309 <211> LENGTH: 41
310 <212> TYPE: DNA
311 <213> ORGANISM: Artificial Sequence
313 <220> FEATURE:
314 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
316 <400> SEQUENCE: 11
317 ggccgctagc ctgcagttat cactctcggg agtgttccca g    41
320 <210> SEQ ID NO: 12
321 <211> LENGTH: 30
322 <212> TYPE: DNA
323 <213> ORGANISM: Artificial Sequence
325 <220> FEATURE:
326 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
328 <400> SEQUENCE: 12
329 ggcggatccg agtggagggc cctgccatcc               30
332 <210> SEQ ID NO: 13

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/462,625

DATE: 05/25/2001

TIME: 14:38:23

Input Set : A:\Pto.amc

Output Set: C:\CRF3\05252001\I462625.raw

L:14 M:270 C: Current Application Number differs, Replaced Application Number

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date

1642

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/462,625

DATE: 05/25/2001

TIME: 13:39:32

Input Set : A:\seqlist- 0652 1630001.txt

Output Set: C:\CRF3\05252001\I462625.raw

Does Not Comply
Corrected Diskette Needed

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4 <110> APPLICANT: Georgiev, Georgii
5      Kiselev, Sergei
6      Prokhorchouk, Egor
7      Ostermann, Elinborg
9 <120> TITLE OF INVENTION: Tumor Growth Inhibition- and Apoptosis-Associated Genes
10     and Polypeptides and Methods of Use Thereof
12 <130> FILE REFERENCE: 0652.1630001
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/462,625
C--> 15 <141> CURRENT FILING DATE: 2000-07-28
17 <150> PRIOR APPLICATION NUMBER: US 08/893,764
18 <151> PRIOR FILING DATE: 1997-07-11
20 <150> PRIOR APPLICATION NUMBER: PCT/EP98/04287
21 <151> PRIOR FILING DATE: 1998-07-10
23 <160> NUMBER OF SEQ ID NOS: 24
25 <170> SOFTWARE: PatentIn Ver. 2.1
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 549
29 <212> TYPE: DNA
30 <213> ORGANISM: Artificial Sequence
32 <220> FEATURE:
33 <221> NAME/KEY: CDS
34 <222> LOCATION: (1)..(549)
36 <220> FEATURE:
37 <223> OTHER INFORMATION: Description of Artificial Sequence: tag7 cDNA
39 <400> SEQUENCE: 1
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41 Met Leu Phe Ala Cys Ala Leu Leu Ala Leu Gly Leu Ala Thr Ser
42 1 5 10 15
44 tgc agt ttc atc gtg ccc cgc agt gag tgg agg gcc ctg cca tcc gag 96
45 Cys Ser Phe Ile Val Pro Arg Ser Glu Trp Arg Ala Leu Pro Ser Glu
46 20 25 30
48 tgc tct agc cgc ctg ggg cac cca gtt cgc tac gtg gtg atc tca cac 144
49 Cys Ser Ser Arg Leu Gly His Pro Val Arg Tyr Val Val Ile Ser His
50 35 40 45
52 aca gcc ggc agc ttc tgc aac agc ccg gac tcc tgt gaa cag cag gcc 192
53 Thr Ala Gly Ser Phe Cys Asn Ser Pro Asp Ser Cys Glu Gln Gln Ala
54 50 55 60
56 cgc aat gtg cag cat tac cac aag aat gag ctg ggc tgg tgc gat gta 240
57 Arg Asn Val Gln His Tyr His Lys Asn Glu Leu Gly Trp Cys Asp Val
58 65 70 75 80
60 gcc tac aac ttc ctt att gga gag gac ggt cat gtc tat gaa ggc cga 288
61 Ala Tyr Asn Phe Leu Ile Gly Glu Asp Gly His Val Tyr Glu Gly Arg
62 85 90 95
64 ggc tgg aac atc aag ggt gac cac aca ggg ccc atc tgg aat ccc atg 336
65 Gly Trp Asn Ile Lys Gly Asp His Thr Gly Pro Ile Trp Asn Pro Met
66 100 105 110
68 tct att ggc atc acc ttc atg ggg aac ttc atg gac cgg gta ccc gca 384

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RAW SEQUENCE LISTING

DATE: 05/25/2001

PATENT APPLICATION: US/09/462,625

TIME: 13:39:32

Input Set : A:\seqlist- 0652 1630001.txt

Output Set: C:\CRF3\05252001\I462625.raw

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72 aag cgg gcc ctc cgt gct gcc cta aat ctt ctg gaa tgt ggg gtg tct 432
73 Lys Arg Ala Leu Arg Ala Ala Leu Asn Leu Leu Glu Cys Gly Val Ser
74      130      135      140
76 cgg ggc ttc ctg aga tcc aac tat gaa gtc aaa gga cac cgg gat gtg 480
77 Arg Gly Phe Leu Arg Ser Asn Tyr Glu Val Lys Gly His Arg Asp Val
78 145      150      155      160
80 caa agc act ctc tct cca ggt gac caa ctc tat cag gtc atc caa agc 528
81 Gln Ser Thr Leu Ser Pro Gly Asp Gln Leu Tyr Gln Val Ile Gln Ser
82      165      170      175
84 tgg gaa cac tac cga gag tga 549
85 Trp Glu His Tyr Arg Glu
86      180
89 <210> SEQ ID NO: 2
90 <211> LENGTH: 182
91 <212> TYPE: PRT
92 <213> ORGANISM: Artificial Sequence
W--> 93 <220> FEATURE: ← add name identifier
93 <223> OTHER INFORMATION: Description of Artificial Sequence: tag7 cDNA
95 <400> SEQUENCE: 2
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97 1 5 10 15
98 Cys Ser Phe Ile Val Pro Arg Ser Glu Trp Arg Ala Leu Pro Ser Glu
99 20 25 30
100 Cys Ser Ser Arg Leu Gly His Pro Val Arg Tyr Val Val Ile Ser His
101 35 40 45
102 Thr Ala Gly Ser Phe Cys Asn Ser Pro Asp Ser Cys Glu Gln Gln Ala
103 50 55 60
104 Arg Asn Val Gln His Tyr His Lys Asn Glu Leu Gly Trp Cys Asp Val
105 65 70 75 80
106 Ala Tyr Asn Phe Leu Ile Gly Glu Asp Gly His Val Tyr Glu Gly Arg
107 85 90 95
108 Gly Trp Asn Ile Lys Gly Asp His Thr Gly Pro Ile Trp Asn Pro Met
109 100 105 110
110 Ser Ile Gly Ile Thr Phe Met Gly Asn Phe Met Asp Arg Val Pro Ala
111 115 120 125
112 Lys Arg Ala Leu Arg Ala Ala Leu Asn Leu Leu Glu Cys Gly Val Ser
113 130 135 140
114 Arg Gly Phe Leu Arg Ser Asn Tyr Glu Val Lys Gly His Arg Asp Val
115 145 150 155 160
116 Gln Ser Thr Leu Ser Pro Gly Asp Gln Leu Tyr Gln Val Ile Gln Ser
117 165 170 175
118 Trp Glu His Tyr Arg Glu
119 180
123 <210> SEQ ID NO: 3
124 <211> LENGTH: 718
125 <212> TYPE: DNA
126 <213> ORGANISM: Homo sapiens

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RAW SEQUENCE LISTING

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TIME: 13:39:32

Input Set : A:\seqlist- 0652 1630001.txt

Output Set: C:\CRF3\05252001\I462625.raw

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128 <220> FEATURE:
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130 <222> LOCATION: (1)..(67)
132 <220> FEATURE:
133 <221> NAME/KEY: CDS
134 <222> LOCATION: (68)..(643)
136 <220> FEATURE:
137 <221> NAME/KEY: 3'UTR
138 <222> LOCATION: (644)..(718)
140 <220> FEATURE:
141 <221> NAME/KEY: polyA_site
142 <222> LOCATION: (712)..(714)
144 <400> SEQUENCE: 3
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147 cgcctct atg ctg ctt gcc tgg gct ctc ccc agc ctc ctt cga ctc gga 109
148 Met Leu Leu Ala Trp Ala Leu Pro Ser Leu Leu Arg Leu Gly
149 1 5 10
151 gcg gct cag gag aca gaa gac ccg gcc tgc tgc agc ccc ata gtg ccc 157
152 Ala Ala Gln Glu Thr Glu Asp Pro Ala Cys Cys Ser Pro Ile Val Pro
153 15 20 25 30
155 cgg aac gag tgg aag gcc ctg gca tca gag tgc gcc cag cac ctg agc 205
156 Arg Asn Glu Trp Lys Ala Leu Ala Ser Glu Cys Ala Gln His Leu Ser
157 35 40 45
159 ctg ccc tta cgc tat gtg gtg gta tgc cac acg gcg ggc agc agc tgc 253
160 Leu Pro Leu Arg Tyr Val Val Val Ser His Thr Ala Gly Ser Ser Cys
161 50 55 60
163 aac acc ccc gcc tgc tgc cag cag cag gcc cgg aat gtg cag cac tac 301
164 Asn Thr Pro Ala Ser Cys Gln Gln Gln Ala Arg Asn Val Gln His Tyr
165 65 70 75
167 cac atg aag aca ctg ggc tgg tgc gac gtg ggc tac aac ttc ctg att 349
168 His Met Lys Thr Leu Gly Trp Cys Asp Val Gly Tyr Asn Phe Leu Ile
169 80 85 90
171 gga gaa gac ggg ctc gta tac gag ggc cgt ggc tgg aac ttc acg ggt 397
172 Gly Glu Asp Gly Leu Val Tyr Glu Gly Arg Gly Trp Asn Phe Thr Gly
173 95 100 105 110
175 gcc cac tca ggt cac tta tgg aac ccc atg tcc att ggc atc agc ttc 445
176 Ala His Ser Gly His Leu Trp Asn Pro Met Ser Ile Gly Ile Ser Phe
177 115 120 125
179 atg ggc aac tac atg gat cgg gtg ccc aca ccc cag gcc atc cgg gca 493
180 Met Gly Asn Tyr Met Asp Arg Val Pro Thr Pro Gln Ala Ile Arg Ala
181 130 135 140
183 gcc cag ggt cta ctg gcc tgc ggt gtg gct cag gga gcc ctg agg tcc 541
184 Ala Gln Gly Leu Leu Ala Cys Gly Val Ala Gln Gly Ala Leu Arg Ser
185 145 150 155
187 aac tat gtg ctc aaa gga cac cgg gat gtg cag cgt aca ctc tct cca 589
188 Asn Tyr Val Leu Lys Gly His Arg Asp Val Gln Arg Thr Leu Ser Pro
189 160 165 170
191 ggc aac cag ctc tac cac ctc atc cag aat tgg cca cac tac cgc tcc 637
192 Gly Asn Gln Leu Tyr His Leu Ile Gln Asn Trp Pro His Tyr Arg Ser

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/462,625

DATE: 05/25/2001

TIME: 13:39:32

Input Set : A:\seqlist- 0652 1630001.txt

Output Set: C:\CRF3\05252001\I462625.raw

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193 175          180          185          190
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196 Pro
198 actgtctcct tctccaataa agatg 718
201 <210> SEQ ID NO: 4
202 <211> LENGTH: 191
203 <212> TYPE: PRT
204 <213> ORGANISM: Homo sapiens
206 <400> SEQUENCE: 4
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209 Gln Glu Thr Glu Asp Pro Ala Cys Cys Ser Pro Ile Val Pro Arg Asn
210 20 25 30
211 Glu Trp Lys Ala Leu Ala Ser Glu Cys Ala Gln His Leu Ser Leu Pro
212 35 40 45
213 Leu Arg Tyr Val Val Val Ser His Thr Ala Gly Ser Ser Cys Asn Thr
214 50 55 60
215 Pro Ala Ser Cys Gln Gln Ala Arg Asn Val Gln His Tyr His Met
216 65 70 75 80
217 Lys Thr Leu Gly Trp Cys Asp Val Gly Tyr Asn Phe Leu Ile Gly Glu
218 85 90 95
219 Asp Gly Leu Val Tyr Glu Gly Arg Gly Trp Asn Phe Thr Gly Ala His
220 100 105 110
221 Ser Gly His Leu Trp Asn Pro Met Ser Ile Gly Ile Ser Phe Met Gly
222 115 120 125
223 Asn Tyr Met Asp Arg Val Pro Thr Pro Gln Ala Ile Arg Ala Ala Gln
224 130 135 140
225 Gly Leu Leu Ala Cys Gly Val Ala Gln Gly Ala Leu Arg Ser Asn Tyr
226 145 150 155 160
227 Val Leu Lys Gly His Arg Asp Val Gln Arg Thr Leu Ser Pro Gly Asn
228 165 170 175
229 Gln Leu Tyr His Leu Ile Gln Asn Trp Pro His Tyr Arg Ser Pro
230 180 185 190
234 <210> SEQ ID NO: 5
235 <211> LENGTH: 14
236 <212> TYPE: DNA
237 <213> ORGANISM: Artificial Sequence
239 <220> FEATURE:
240 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
242 <400> SEQUENCE: 5
243 tttttttttt ttac 14
246 <210> SEQ ID NO: 6
247 <211> LENGTH: 10
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial Sequence
251 <220> FEATURE:
252 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
254 <400> SEQUENCE: 6
255 aatcgggctg 10

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/462,625

DATE: 05/25/2001

TIME: 13:39:32

Input Set : A:\seqlist- 0652 1630001.txt

Output Set: C:\CRF3\05252001\I462625.raw

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258 <210> SEQ ID NO: 7
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260 <212> TYPE: DNA
261 <213> ORGANISM: Artificial Sequence
263 <220> FEATURE:
264 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
266 <400> SEQUENCE: 7
267 agtcagccac                                     10
270 <210> SEQ ID NO: 8
271 <211> LENGTH: 18
272 <212> TYPE: DNA
273 <213> ORGANISM: Artificial Sequence
275 <220> FEATURE:
276 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
278 <400> SEQUENCE: 8
279 gaaagggcct cgtgatac                             18
282 <210> SEQ ID NO: 9
283 <211> LENGTH: 36
284 <212> TYPE: DNA
285 <213> ORGANISM: Artificial Sequence
287 <220> FEATURE:
288 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
290 <400> SEQUENCE: 9
291 cagggccctc cactcacttg cataggcatt tgtagc         36
294 <210> SEQ ID NO: 10
295 <211> LENGTH: 36
296 <212> TYPE: DNA
297 <213> ORGANISM: Artificial Sequence
299 <220> FEATURE:
300 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
302 <400> SEQUENCE: 10
303 gctacaaatg cctatgcaag tgagtggagg gccctg         36
306 <210> SEQ ID NO: 11
307 <211> LENGTH: 41
308 <212> TYPE: DNA
309 <213> ORGANISM: Artificial Sequence
311 <220> FEATURE:
312 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
314 <400> SEQUENCE: 11
315 ggccgctagc ctgcagttat cactctcggt agtgttccca g    41
318 <210> SEQ ID NO: 12
319 <211> LENGTH: 30
320 <212> TYPE: DNA
321 <213> ORGANISM: Artificial Sequence
323 <220> FEATURE:
324 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Primer
326 <400> SEQUENCE: 12
327 ggcggatccg agtggagggc cctgccatcc               30
330 <210> SEQ ID NO: 13

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/462,625

DATE: 05/25/2001

TIME: 13:39:33

Input Set : A:\seqlist- 0652 1630001.txt

Output Set: C:\CRF3\05252001\I462625.raw

L:14 M:270 C: Current Application Number differs, Replaced Application Number

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:93 M:258 W: Mandatory Feature missing, <220> FEATURE: